

SEQUENCE LISTING

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<110> AFEYAN, NOUBAR B.
        LEE, FRANK D.
       WONG, GORDON G.
       DAS GUPTA, RUCHIRA
       BAYNES, BRIAN
 <120> ADZYMES AND USES THEREOF
 <130> COTH-P02-001
 <140> 10/650,591
 <141> 2003-08-27
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 <150> 60/423,754
 <151> 2002-11-05
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 Glu Glu Thr Ala Arg Phe Gln Pro Gly Tyr Arg Ser
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 <400> 2
 Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
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      peptide
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Asp Tyr Lys Asp Asp Asp Lys
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Tyr Pro Tyr Asp Val Pro Asp Tyr Ala
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      peptide
Glu Asp Gln Val Asp Pro Arg Leu Ile Asp Gly Lys
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      peptide
<400> 6
Tyr Thr Asp Ile Glu Met Asn Arg Leu Gly Lys
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<212> PRT
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<400> 7
Ser Ser Ser Gly
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<400> 8
Ser Gly Gly Gly
                5
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      peptide
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<221> MOD RES
<222> (3)..(4)
<223> Any amino acid
<400> 9
His Glu Xaa Xaa His
<210> 10
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      6x His tag
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His His His His His
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1 5
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<210> 11
<211> 26
<212> PRT
<213> Human immunodeficiency virus 1
Cys Phe Ile Thr Lys Ala Leu Gly Ile Ser Tyr Gly Arg Lys Lys Arg
                5
                                     10
Arg Gln Arg Arg Pro Pro Gln Gly Ser
            20
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<213> Artificial Sequence
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Cys Met His Ile Glu Ser Leu Asp Ser Tyr Thr Cys
<210> 13
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<213> Artificial Sequence
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Cys Met Tyr Ile Glu Ala Leu Asp Lys Tyr Ala Cys
                5
<210> 14
<211> 32
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      peptide
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<221> MOD RES
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<222> (1)..(1)
<223> A unique residue, such as cysteine or lysine, that
      facilitates chemical conjugation of the internalizing
      peptide to a targeting protein conjugate
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<221> MOD RES
<222> (2)..(3)
<223> Any residues selected to modulate the affinity of the
      internalizing peptide for different membranes
<220>
<223> see specification as filed for detailed description of
      substitutions and preferred embodiments
<400> 14
Xaa Xaa Xaa Glu Ala Ala Leu Ala Glu Ala Leu Ala Glu Ala Leu Ala
                                    10
Glu Ala Leu Ala Glu Ala Leu Ala Leu Glu Ala Leu Ala Ala
            20
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      peptide
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Ala Leu Trp His Trp Trp His
<210> 16
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<222> (1)..(1)
<223> Thr or Ser
<400> 16
Xaa Trp Leu His Trp Trp Ala
                5
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<210> 17
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<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 17
Gly Gly Gly Ser
                5
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<211> 6
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<213> Influenza A virus
<400> 18
Asp Val Pro Asp Tyr Ala
<210> 19
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Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser
                5
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<210> 20
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Gly Gly Val Arg
<210> 21
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<223> Description of Artificial Sequence: Synthetic
 polypeptide

<400> 21

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Ala Ala Gln Pro Ala Arg Arg Ala Val Arg Ser 20 25 30

Leu Met Thr Ala Thr Ser Glu Tyr Gln Thr Phe Phe Asn Pro Arg Thr 35 40 45

Phe Gly Ser Gly Glu Ala Asp Cys Gly Leu Arg Pro Leu Phe Glu Lys 50 55 60

Lys Ser Leu Glu Asp Lys Thr Glu Arg Glu Leu Leu Glu Ser Tyr Ile 65 70 75 80

Asp Gly Arg Ile Val Glu Gly Ser Asp Ala Glu Ile Gly Met Ser Pro 85 90 95

Trp Gln Val Met Leu Phe Arg Lys Ser Pro Gln Glu Leu Leu Cys Gly 100 105 110

Ala Ser Leu Ile Ser Asp Arg Trp Val Leu Thr Ala Ala His Cys Leu 115 120 125

Leu Tyr Pro Pro Trp Asp Lys Asn Phe Thr Glu Asn Asp Leu Leu Val

Arg Ile Gly Lys His Ser Arg Thr Arg Tyr Glu Arg Asn Ile Glu Lys 145 150 155 160

Ile Ser Met Leu Glu Lys Ile Tyr Ile His Pro Arg Tyr Asn Trp Arg 165 170 175

Glu Asn Leu Asp Arg Asp Ile Ala Leu Met Lys Leu Lys Lys Pro Val 180 185 190

Ala Phe Ser Asp Tyr Ile His Pro Val Cys Leu Pro Asp Arg Glu Thr
195 200 205

Ala	Ala 210	Ser	Leu	Leu	Gln	Ala 215	Gly	Tyr	Lys	Gly	Arg 220	Val	Thr	Gly	Trp
Gly 225	Asn	Leu	Lys	Glu	Thr 230	Trp	Thr	Ala	Asn	Val 235	Gly	Lys	Gly	Gln	Pro 240
Ser	Val	Leu	Gln	Val 245	Val	Asn	Leu	Pro	Ile 250	Val	Glu	Arg	Pro	Val 255	Cys
Lys	Asp	Ser	Thr 260	Arg	Ile	Arg	Ile	Thr 265	Asp	Asn	Met	Phe	Cys 270	Ala	Gly
Tyr	Lys	Pro 275	Asp	Glu	Gly	Lys	Arg 280	Gly	Asp	Ala	Cys	Glu 285	Gly	Asp	Ser
Gly	Gly 290	Pro	Phe	Val	Met	Lys 295	Ser	Pro	Phe	Asn	Asn 300	Arg	Trp	Tyr	Gln
Met 305	Gly	Ile	Val	Ser	Trp 310	Gly	Glu	Gly	Cys	Asp 315	Arg	Asp	Gly	Lys	Tyr 320
Gly	Phe	Tyr	Thr	His 325	Val	Phe	Arg	Leu	Lys 330	Lys	Trp	Ile	Gln	Lys 335	Val
Ile	Asp	Gln	Phe 340	Gly	Glu	Gly	Gly	Gly 345	Gly	Ser	Gly	Gly	Gly 350	Gly	Ser
Gly	Gly	Gly 355	Gly	Ser	Met	Glu	Val 360	Gln	Leu	Leu	Glu	Ser 365	Gly	Gly	Asp
Leu	Val 370	Lys	Pro	Gly	Gly	Ser 375	Leu	Lys	Leu	Ser	Cys 380	Ala	Ala	Ser	Gly
Phe 385	Thr	Phe	Ser	Thr	Tyr 390	Gly	Met	Ser	Trp	Val 395	Arg	Gln	Thr	Pro	Asp 400
Lys	Arg	Leu	Glu	Trp 405	Val	Ala	Thr	Ile	Ser 410	Asn	Gly	Gly	Gly	Tyr 415	Thr
Tyr	Tyr	Pro	Asp 420	Ser	Val	Lys	Gly	Arg 425	Phe	Thr	Ile	Ser	Arg 430	Asp	Asn

Ala Lys Asn Thr Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser Glu Asp 435 440 445

Thr Ala Met Tyr Tyr Cys Ala Arg Arg Glu Arg Tyr Asp Glu Asn Gly 450 455 460

Phe Ala Tyr Trp Gly Arg Gly Thr Leu Val Thr Val Ser Ala Gly Gly 465 470 475 480

Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Asp Ile Val 485 490 495

Met Ser Gln Ser Pro Ser Ser Leu Ala Val Ser Val Gly Glu Lys Ile 500 505 510

Thr Met Ser Cys Lys Ser Ser Gln Ser Leu Phe Asn Ser Gly Lys Gln 515 520 525

Lys Asn Tyr Leu Thr Trp Tyr Gln Gln Lys Pro Gly Gln Ser Pro Lys 530 540

Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Asp Arg 545 550 555 560

Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser 565 570 575

Val Lys Ala Glu Asp Leu Ala Val Tyr Tyr Cys Gln Asn Asp Tyr Ser 580 585 590

His Pro Leu Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg Ala 595 600 605

Asp Ala Ala Pro Thr Ala Arg Gly Gly Pro Glu Gln Lys Leu Ile Ser 610 620

Glu Glu Asp Leu Asn Ser Ala Val Asp His His His His His 625 630 635

<210> 22

<211> 639

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 22

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Ala Ala Gln Pro Ala Arg Arg Ala Val Arg Ser 20 25 30

Leu Met Glu Val Gln Leu Leu Glu Ser Gly Gly Asp Leu Val Lys Pro 35 40 45

Gly Gly Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser 50 55 60

Thr Tyr Gly Met Ser Trp Val Arg Gln Thr Pro Asp Lys Arg Leu Glu 65 70 75 80

Trp Val Ala Thr Ile Ser Asn Gly Gly Gly Tyr Thr Tyr Tyr Pro Asp 85 90 95

Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ala Lys Asn Thr 100 105 110

Leu Tyr Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Met Tyr 115 120 125

Tyr Cys Ala Arg Arg Glu Arg Tyr Asp Glu Asn Gly Phe Ala Tyr Trp 130 135 140

Gly Arg Gly Thr Leu Val Thr Val Ser Ala Gly Gly Gly Gly Ser Gly 145 150 155 160

Gly Gly Gly Ser Gly Gly Gly Ser Asp Ile Val Met Ser Gln Ser 165 170 175

Pro Ser Ser Leu Ala Val Ser Val Gly Glu Lys Ile Thr Met Ser Cys 180 185 190

Lys Ser Ser Gln Ser Leu Phe Asn Ser Gly Lys Gln Lys Asn Tyr Leu 195 200 205

Thr Trp Tyr 210	Gln Gln I	Lys Pro 215	Gly	Gln	Ser	Pro	Lys 220	Leu	Leu	Ile	Tyr
Trp Ala Ser 225	_	Glu Ser 230	Gly	Val	Pro	Asp 235	Arg	Phe	Thr	Gly	Ser 240
Gly Ser Gly	Thr Asp I	Phe Thr	Leu '		Ile 250	Ser	Ser	Val	Lys	Ala 255	Glu
Asp Leu Ala	Val Tyr 1 260	Tyr Cys		Asn 265	Asp	Tyr	Ser	His	Pro 270	Leu	Thr
Phe Gly Gly 275	-	Lys Leu	Glu 280	Ile	Lys	Arg	Ala	Asp 285	Ala	Ala	Pro
Thr Gly Gly 290	Gly Gly S	Ser Gly 295	Gly	Gly	Gly	Ser	Gly 300	Gly	Gly	Gly	Ser
Met Thr Ala		Glu Tyr 310	Gln '	Thr	Phe	Phe 315	Asn	Pro	Arg	Thr	Phe 320
Gly Ser Gly	Glu Ala A 325	Asp Cys	Gly :		Arg 330	Pro	Leu	Phe	Glu	Lys 335	Lys
Ser Leu Glu	Asp Lys 3	Thr Glu	_	Glu : 345	Leu	Leu	Glu	Ser	Tyr 350	Ile	Asp
Gly Arg Ile 355		Gly Ser	Asp 3	Ala	Glu	Ile	Gly	Met 365	Ser	Pro	Trp
Gln Val Met 370	Leu Phe I	Arg Lys 375	Ser :	Pro (Gln	Glu	Leu 380	Leu	Cys	Gly	Ala
Ser Leu Ile 385		Arg Trp 390	Val :	Leu '		Ala 395	Ala	His	Cys	Leu	Leu 400
Tyr Pro Pro	Trp Asp I 405	Lys Asn	Phe '		Glu 410	Asn	Asp	Leu	Leu	Val 415	Arg
Ile Gly Lys	His Ser A	Arg Thr	_	Tyr (425	Glu	Arg	Asn	Ile	Glu 430	Lys	Ile
Ser Met Leu	Glu Lys I	Ile Tyr	Ile I	His 1	Pro	Arg	Tyr	Asn	Trp	Arg	Glu

435 440 445

Asn Leu Asp Arg Asp Ile Ala Leu Met Lys Leu Lys Lys Pro Val Ala 450 455 460

Phe Ser Asp Tyr Ile His Pro Val Cys Leu Pro Asp Arg Glu Thr Ala 465 470 475 480

Ala Ser Leu Leu Gln Ala Gly Tyr Lys Gly Arg Val Thr Gly Trp Gly
485 490 495

Asn Leu Lys Glu Thr Trp Thr Ala Asn Val Gly Lys Gly Gln Pro Ser 500 505 510

Val Leu Gln Val Val Asn Leu Pro Ile Val Glu Arg Pro Val Cys Lys 515 520 525

Asp Ser Thr Arg Ile Arg Ile Thr Asp Asn Met Phe Cys Ala Gly Tyr 530 535 540

Lys Pro Asp Glu Gly Lys Arg Gly Asp Ala Cys Glu Gly Asp Ser Gly 545 550 555 560

Gly Pro Phe Val Met Lys Ser Pro Phe Asn Asn Arg Trp Tyr Gln Met 565 570 575

Gly Ile Val Ser Trp Gly Glu Gly Cys Asp Arg Asp Gly Lys Tyr Gly
580 585 590

Phe Tyr Thr His Val Phe Arg Leu Lys Lys Trp Ile Gln Lys Val Ile
595 600 605

Asp Gln Phe Gly Glu Ala Arg Gly Gly Pro Glu Gln Lys Leu Ile Ser 610 620

Glu Glu Asp Leu Asn Ser Ala Val Asp His His His His His 625 630 635

<210> 23

<211> 30

<212> DNA

<213> Artificial Sequence

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<210><211><212><213>	31	
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<400> acgcco	24 cctcg agcagttggt gcagcatcag c	31
<210><211><211><212><213>	33	
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<400> cccgga	25 aagct taatgaccgc caccagtgag tac	33
<210><211><211><212><213>	31	
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<400> ggccc	26 etcga gcctctccaa actgatcaat g	31
<210><211><212><212><213>	72	
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<400>	27 agagg gaggcggtgg gtctggtggg ggcggtagtg gcggaggtgg gagcatggag	60

gtgcag	gctgt tg	72
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	ccatg ctcccacctc cgccactacc gccccacca gacccaccgc ctccctctcc	60 73
aaacc	gatca atg	/ 3
<210><211><211><212><213>	75	
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<400> gcacca	29 aactg gaggeggtgg gtetggtggg ggeggtagtg geggaggtgg gageatgace	60
gccaco	cagtg agtac	75
<210><211><211><212><213>	75	
<220> <223>	Description of Artificial Sequence: Synthetic oligonucleotide	
<400> ggtggd	30 eggte atgeteccae etecgecaet acegececca ceagacecae egectecagt	60
tggtgd	cagca tcagc	75
<210><211><211><212><213>	33	
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Tyr Pro Tyr Asp Val Pro Asp Tyr Ala Ser Gly Ser Gly Ser Ser Gly
Ser Gly Ser Ser Gly Ser Gly Ser Gly Ser Gly Ser Gly Val
Arg
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<212> PRT
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<400> 32
Ile Thr Pro Arg
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<210> 33
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<400> 33
Ile Thr Leu Arg
<210> 34
<211> 18
<212> PRT
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Tyr Pro Tyr Asp Val Pro Asp Tyr Ala Gly Ser Gly Asp Tyr Lys Ala
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Phe Asp

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<223> Description of Artificial Sequence: Synthetic
     peptide
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Gly Gly Gly Ser Gly Gly Gly Ser
               5
<210> 36
<211> 20
<212> PRT
<213> Artificial Sequence
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     peptide
<400> 36
Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser Gly
Gly Gly Gly Ser
           20
<210> 37
<211> 30
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 37
Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser Gly
Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser
                               25
<210> 38
<211> 40
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 38
Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser Gly
Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly
Gly Gly Ser Gly Gly Gly Ser
       35
<210> 39
<211> 50
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
     peptide
Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly
               5
                                  10
                                                      15
Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly
Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly Ser Gly Gly Gly
Gly Ser
   50
<210> 40
<211> 280
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<223> Description of Artificial Sequence: Synthetic
     polypeptide
<400> 40
Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro
               5
                                  10
                                                      15
```

- Gly Ser Thr Gly Asp Ile Ala Pro Phe Asp Asp Asp Asp Lys Ile Val 20 25 30
- Gly Gly Tyr Asn Cys Glu Glu Asn Ser Val Pro Tyr Gln Val Ser Leu 35 40 45
- Asn Ser Gly Tyr His Phe Cys Gly Gly Ser Leu Ile Asn Glu Gln Trp 50 60
- Val Val Ser Ala Gly His Cys Tyr Lys Ser Arg Ile Gln Val Arg Leu 65 70 75 80
- Gly Glu His Asn Ile Glu Val Leu Glu Gly Asn Glu Gln Phe Ile Asn 85 90 95
- Ala Ala Lys Ile Ile Arg His Pro Gln Tyr Asp Arg Lys Thr Leu Asn 100 105 110
- Asn Asp Ile Met Leu Ile Lys Leu Ser Ser Arg Ala Val Ile Asn Ala 115 120 125
- Arg Val Ser Thr Ile Ser Leu Pro Thr Ala Pro Pro Ala Thr Gly Thr 130 135 140
- Lys Cys Leu Ile Ser Gly Trp Gly Asn Thr Ala Ser Ser Gly Ala Asp 145 150 155 160
- Tyr Pro Asp Glu Leu Gln Cys Leu Asp Ala Pro Val Leu Ser Gln Ala 165 170 175
- Lys Cys Glu Ala Ser Tyr Pro Gly Lys Ile Thr Ser Asn Met Phe Cys 180 185 190
- Val Gly Phe Leu Glu Gly Gly Lys Asp Ser Cys Gln Gly Asp Ser Gly 195 200 205
- Gly Pro Val Val Cys Asn Gly Gln Leu Gln Gly Val Val Ser Trp Gly 210 215 220
- Asp Gly Cys Ala Gln Lys Asn Lys Pro Gly Val Tyr Thr Lys Val Tyr 225 230 235 240

Asn Tyr Val Lys Trp Ile Lys Asn Thr Ile Ala Ala Asn Ser Thr Arg 245 250 255

Gly Gly Pro Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Ser Ala 260 265 270

Val Asp His His His His His 275 280

<210> 41

<211> 461

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 41

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Ile Ala Pro Phe Asp Asp Asp Lys Ile Val 20 25 30

Gly Gly Tyr Asn Cys Glu Glu Asn Ser Val Pro Tyr Gln Val Ser Leu 35 40 45

Asn Ser Gly Tyr His Phe Cys Gly Gly Ser Leu Ile Asn Glu Gln Trp 50 55 60

Val Val Ser Ala Gly His Cys Tyr Lys Ser Arg Ile Gln Val Arg Leu 65 70 75 80

Gly Glu His Asn Ile Glu Val Leu Glu Gly Asn Glu Gln Phe Ile Asn 85 90 95

Ala Ala Lys Ile Ile Arg His Pro Gln Tyr Asp Arg Lys Thr Leu Asn 100 105 110

Asn Asp Ile Met Leu Ile Lys Leu Ser Ser Arg Ala Val Ile Asn Ala 115 120 125

Arg Val Ser Thr Ile Ser Leu Pro Thr Ala Pro Pro Ala Thr Gly Thr 130 135 140

Ly:	s Cys	Leu	Ile	Ser	Gly 150	Trp	Gly	Asn	Thr	Ala 155	Ser	Ser	Gly	Ala	Asp 160
Ту	r Pro	Asp	Glu	Leu 165	Gln	Cys	Leu	Asp	Ala 170	Pro	Val	Leu	Ser	Gln 175	Ala
Lу	s Cys	Glu	Ala 180	Ser	Tyr	Pro	Gly	Lys 185	Ile	Thr	Ser	Asn	Met 190	Phe	Cys
Va:	l Gly	Phe 195		Glu	Gly	Gly	Lys 200	Asp	Ser	Cys	Gln	Gly 205	Asp	Ser	Gly
Gl	y Pro 210		Val	Cys	Asn	Gly 215	Gln	Leu	Gln	Gly	Val 220	Val	Ser	Trp	Gly
As _]	Gly	Cys	Ala	Gln	Lys 230	Asn	Lys	Pro	Gly	Val 235	Tyr	Thr	Lys	Val	Tyr 240
Ası	ı Tyr	Val	Lys	Trp 245	Ile	Lys	Asn	Thr	Ile 250	Ala	Ala	Asn	Ser	Leu 255	Val
Pro) His	Leu	Gly 260	Asp	Arg	Glu	Lys	Arg 265	Asp	Ser	Val	Cys	Pro 270	Gln	Gly
Ly	5 Tyr	Ile 275	His	Pro	Gln	Asn	Asn 280	Ser	Ile	Сув	Cys	Thr 285	Lys	Cys	His
Lу	3 Gly 290		Tyr	Leu	Tyr	Asn 295	Asp	Cys	Pro	Gly	Pro 300	Gly	Gln	Asp	Thr
As _]	Cys	Arg	Glu	Cys	Glu 310	Ser	Gly	Ser	Phe	Thr 315	Ala	Ser	Glu	Asn	His 320
Le	ı Arg	His	Cys	Leu 325	Ser	Cys	Ser	Lys	Cys 330	Arg	Lys	Glu	Met	Gly 335	Gln
Va:	l Glu	Ile	Ser 340	Ser	Сув	Thr	Val	Asp 345	Arg	Asp	Thr	Val	Cys 350	Gly	Cys
Arg	J Lys	Asn 355	Gln	Tyr	Arg	His	Tyr 360	Trp	Ser	Glu	Asn	Leu 365	Phe	Gln	Сув

Phe Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu Ser Cys Gln 370 380

Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg 385 390 395 400

Glu Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Ser Leu Glu Cys 405 410 415

Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly Thr Glu Asp 420 425 430.

Ser Gly Thr Thr Arg Gly Gly Pro Glu Gln Lys Leu Ile Ser Glu Glu
435 440 445

Asp Leu Asn Ser Ala Val Asp His His His His His 450 455 460

<210> 42

<211> 464

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 polypeptide

<400> 42

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Ile Ala Pro Phe Asp Asp Asp Asp Lys Ile Val 20 25 30

Gly Gly Tyr Asn Cys Glu Glu Asn Ser Val Pro Tyr Gln Val Ser Leu 35 40 45

Asn Ser Gly Tyr His Phe Cys Gly Gly Ser Leu Ile Asn Glu Gln Trp 50 55 60

Val Val Ser Ala Gly His Cys Tyr Lys Ser Arg Ile Gln Val Arg Leu 65 70 75 80

Gly Glu His Asn Ile Glu Val Leu Glu Gly Asn Glu Gln Phe Ile Asn 85 90 95

- Ala Ala Lys Ile Ile Arg His Pro Gln Tyr Asp Arg Lys Thr Leu Asn 100 105 110
- Asn Asp Ile Met Leu Ile Lys Leu Ser Ser Arg Ala Val Ile Asn Ala 115 120 125
- Arg Val Ser Thr Ile Ser Leu Pro Thr Ala Pro Pro Ala Thr Gly Thr 130 135 140
- Lys Cys Leu Ile Ser Gly Trp Gly Asn Thr Ala Ser Ser Gly Ala Asp 145 150 155 160
- Tyr Pro Asp Glu Leu Gln Cys Leu Asp Ala Pro Val Leu Ser Gln Ala 165 170 175
- Lys Cys Glu Ala Ser Tyr Pro Gly Lys Ile Thr Ser Asn Met Phe Cys 180 185 190
- Val Gly Phe Leu Glu Gly Gly Lys Asp Ser Cys Gln Gly Asp Ser Gly
 195 200 205
- Gly Pro Val Val Cys Asn Gly Gln Leu Gln Gly Val Val Ser Trp Gly 210 215 220
- Asp Gly Cys Ala Gln Lys Asn Lys Pro Gly Val Tyr Thr Lys Val Tyr 225 230 235 240
- Asn Tyr Val Lys Trp Ile Lys Asn Thr Ile Ala Ala Asn Ser Ala Ala 245 250 255
- Ala Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg Asp Ser Val Cys 260 265 270
- Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr 275 280 285
- Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro Gly 290 295 300
- Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser 305 310 315 320

Glu Asn His Leu Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys Glu 325 330 335

Met Gly Gln Val Glu Ile Ser Ser Cys Thr Val Asp Arg Asp Thr Val 340 345 350

Cys Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn Leu 355 360 365

Phe Gln Cys Phe Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu 370 380

Ser Cys Gln Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe 385 390 395 400

Phe Leu Arg Glu Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser 405 410 415

Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly
420 425 430

Thr Glu Asp Ser Gly Thr Thr Arg Gly Gly Pro Glu Gln Lys Leu Ile 435 440 445

Ser Glu Glu Asp Leu Asn Ser Ala Val Asp His His His His His 450 455 460

<210> 43

<211> 485

<212> PRT

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic polypeptide

<400> 43

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro 1 5 10 15

Gly Ser Thr Gly Asp Ile Ala Pro Phe Asp Asp Asp Lys Ile Val 20 25 30

Gly Gly Tyr Asn Cys Glu Glu Asn Ser Val Pro Tyr Gln Val Ser Leu 35 40 45

- Asn Ser Gly Tyr His Phe Cys Gly Gly Ser Leu Ile Asn Glu Gln Trp 50 55 60
- Val Val Ser Ala Gly His Cys Tyr Lys Ser Arg Ile Gln Val Arg Leu 65 70 75 80
- Gly Glu His Asn Ile Glu Val Leu Glu Gly Asn Glu Gln Phe Ile Asn 85 90 95
- Ala Ala Lys Ile Ile Arg His Pro Gln Tyr Asp Arg Lys Thr Leu Asn 100 105 110
- Asn Asp Ile Met Leu Ile Lys Leu Ser Ser Arg Ala Val Ile Asn Ala 115 120 125
- Arg Val Ser Thr Ile Ser Leu Pro Thr Ala Pro Pro Ala Thr Gly Thr 130 135 140
- Lys Cys Leu Ile Ser Gly Trp Gly Asn Thr Ala Ser Ser Gly Ala Asp 145 150 155 160
- Tyr Pro Asp Glu Leu Gln Cys Leu Asp Ala Pro Val Leu Ser Gln Ala 165 170 175
- Lys Cys Glu Ala Ser Tyr Pro Gly Lys Ile Thr Ser Asn Met Phe Cys 180 185 190
- Val Gly Phe Leu Glu Gly Gly Lys Asp Ser Cys Gln Gly Asp Ser Gly 195 200 205
- Gly Pro Val Val Cys Asn Gly Gln Leu Gln Gly Val Val Ser Trp Gly 210 220
- Asp Gly Cys Ala Gln Lys Asn Lys Pro Gly Val Tyr Thr Lys Val Tyr 225 230 235 240
- Asn Tyr Val Lys Trp Ile Lys Asn Thr Ile Ala Ala Asn Ser Ala Ala 245 250 255
- Ala Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Ser 260 265 270

Gly Gly Gly Ser Arg Leu Val Pro His Leu Gly Asp Arg Glu Lys 275 280 285

Arg Asp Ser Val Cys Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn 290 295 300

Ser Ile Cys Cys Thr Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp 305 310 315 320

Cys Pro Gly Pro Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly 325 330 335

Ser Phe Thr Ala Ser Glu Asn His Leu Arg His Cys Leu Ser Cys Ser 340 345 350

Lys Cys Arg Lys Glu Met Gly Gln Val Glu Ile Ser Ser Cys Thr Val 355 360 365

Asp Arg Asp Thr Val Cys Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr 370 375 380

Trp Ser Glu Asn Leu Phe Gln Cys Phe Asn Cys Ser Leu Cys Leu Asn 385 390 395 400

Gly Thr Val His Leu Ser Cys Gln Glu Lys Gln Asn Thr Val Cys Thr 405 410 415

Cys His Ala Gly Phe Phe Leu Arg Glu Asn Glu Cys Val Ser Cys Ser 420 425 430

Asn Cys Lys Lys Ser Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile 435 440 445

Glu Asn Val Lys Gly Thr Glu Asp Ser Gly Thr Thr Arg Gly Gly Pro 450 455 460

Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Ser Ala Val Asp His 465 470 475 480

His His His His

<211> 239

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic
 polypeptide

<400> 44

Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro

1 10 15

Gly Ser Thr Gly Asp Ala Ala Gln Pro Ala Arg Arg Ala Val Arg Ser 20 25 30

Leu Val Pro His Leu Gly Asp Arg Glu Lys Arg Asp Ser Val Cys Pro 35 40 45

Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr Lys 50 55 60

Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro Gly Gln 65 70 75 80

Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser Glu 85 90 95

Asn His Leu Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys Glu Met 100 105 110

Gly Gln Val Glu Ile Ser Ser Cys Thr Val Asp Arg Asp Thr Val Cys 115 120 125

Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn Leu Phe 130 135

Gln Cys Phe Asn Cys Ser Leu Cys Leu Asn Gly Thr Val His Leu Ser 145 150 155 160

Cys Gln Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe 165 170 175

Leu Arg Glu Asn Glu Cys Val Ser Cys Ser Asn Cys Lys Lys Ser Leu 180 185 190 Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly Thr 195 200 205

Glu Asp Ser Gly Thr Thr Arg Gly Gly Pro Glu Gln Lys Leu Ile Ser 210 215 220

Glu Glu Asp Leu Asn Ser Ala Val Asp His His His His His E25 230 235

<210> 45

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Synthetic peptide

<400> 45

Arg Pro Leu Ala Leu Trp Arg Ser